5

	Application No.	Applicant(s)					
	10/631,958	KOSSIDA ET AL.					
Notice of Allowability	Examiner	Art Unit					
	Maryam Monshipouri	1653					
The MAILING DATE of this communication appears on the cover sheet with the correspondence address All claims being allowable, PROSECUTION ON THE MERITS IS (OR REMAINS) CLOSED in this application. If not included herewith (or previously mailed), a Notice of Allowance (PTOL-85) or other appropriate communication will be mailed in due course. THIS NOTICE OF ALLOWABILITY IS NOT A GRANT OF PATENT RIGHTS. This application is subject to withdrawal from issue at the initiative of the Office or upon petition by the applicant. See 37 CFR 1.313 and MPEP 1308.							
1. This communication is responsive to							
2. X The allowed claim(s) is/are 10,12,13,14 and 23.							
 3. Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f). a) All b) Some* c) None of the: 1. Certified copies of the priority documents have been received. 2. Certified copies of the priority documents have been received in Application No. 3. Copies of the certified copies of the priority documents have been received in this national stage application from the International Bureau (PCT Rule 17.2(a)). * Certified copies not received: Applicant has THREE MONTHS FROM THE "MAILING DATE" of this communication to file a reply complying with the requirements noted below. Failure to timely comply will result in ABANDONMENT of this application. 							
THIS THREE-MONTH PERIOD IS NOT EXTENDABLE. 4. A SUBSTITUTE OATH OR DECLARATION must be submitted. Note the attached EXAMINER'S AMENDMENT or NOTICE OF INFORMAL PATENT APPLICATION (PTO-152) which gives reason(s) why the oath or declaration is deficient.							
 5. CORRECTED DRAWINGS (as "replacement sheets") must be submitted. (a) including changes required by the Notice of Draftsperson's Patent Drawing Review (PTO-948) attached 1) hereto or 2) to Paper No./Mail Date (b) including changes required by the attached Examiner's Amendment / Comment or in the Office action of Paper No./Mail Date Identifying indicia such as the application number (see 37 CFR 1.84(c)) should be written on the drawings in the front (not the back) of each sheet. Replacement sheet(s) should be labeled as such in the header according to 37 CFR 1.121(d). 							
6. DEPOSIT OF and/or INFORMATION about the deposit of BIOLOGICAL MATERIAL must be submitted. Note the attached Examiner's comment regarding REQUIREMENT FOR THE DEPOSIT OF BIOLOGICAL MATERIAL.							
 Attachment(s) 1. ☑ Notice of References Cited (PTO-892) 2. ☐ Notice of Draftperson's Patent Drawing Review (PTO-948) 3. ☑ Information Disclosure Statements (PTO-1449 or PTO/SB/0 Paper No./Mail Date filed 8/1/03) 4. ☐ Examiner's Comment Regarding Requirement for Deposit of Biological Material 	5. ☐ Notice of Informal P 6. ☑ Interview Summary Paper No./Mail Dat 7. ☑ Examiner's Amendr 8. ☑ Examiner's Stateme 9. ☑ Other <u>See Continua</u>	(PTO-413), te <u>9/14/05</u> . ment/Comment ent of Reasons for Allo					

Continuation Sheet (PTOL-37)

Application No. 10/631,958

Continuation of Attachment(s) 9. Other: see atachments to interview summary.

Application/Control Number: 10/631,958

Art Unit: 1653

An **Examiner's Amendment** to the record appears below. Should the changes and/or additions be unacceptable to applicant, an amendment may be filed as provided by 37 CFR 1.312. To ensure consideration of such an amendment, it MUST be submitted no later than the payment of the issue fee.

Authorization for this Examiner's Amendment was given in a telephone interview with Ms. Lisa M. Hemmendinger, on 9/15/2005.

Examiner's Amendment to the Claims

Cancel claims 11 and 15.

In claim 10, line 2, after "consisting of " delete "(a) ".

In claim 10, line 2, after "SEQ ID NOS:", delete "2".

In claim 10, lines 2-3, delete "and (b) biologically active variants thereof ".

In claim 14, line 2, after "consisting of " delete "(a) ".

In claim 14, line 3, after "SEQ ID NOS:", delete "2".

In claim 14, after "10,", delete "or", and substitute therefor --- and ---.

In claim 14, line 3, delete "and (b) biologically active variants thereof".

In claim 23, line 3, after "SEQ ID NOS:", delete "2".

In claim 23, line 2, after "consisting of " delete "(a) ".

In claim 23, line 3, after "11", delete ", and (b) biologically active variants thereof".

In claim 23, line 6, delete " claim 21' and substitute therefor --- detecting said

polypeptide comprising the steps of: contacting a biological sample with said

antibody to form a reagent -polypeptide complex; and

Application/Control Number: 10/631,958

Art Unit: 1653

detecting the antibody-polypeptide complex. ---.

This application is in condition for allowance except for the presence of claims 1-9, 16-22, 24-77 drawn to non-elected without traverse. Accordingly, claims 1-9, 16-22, 24-77 have been cancelled.

Cancel claims 1-9, 16-22, 24-77.

Examiner's Amendments to the Specification

In page 1, line 2, after "October 4, 2001,", insert --- now abandoned ---.

The following is an **Examiner's Statement of Reasons for Allowance**:

Claims 10, 12-14 and 23 are directed to an isolated ceramide kinase of specific amino acid sequence and a specifically claimed fragment thereof with said activity, fusion proteins and kits comprising said polypepitdes.

Claimed ceramide kinase and said fragment thereof are free of prior art. Further the prior art does not teach or suggest preparing such specifically claimed products.

Hence said products are also novel and non-obvious.

Since said polypepitdes are both novel and non-obvious fusion proteins and kits comprising said polypeptides are also novel and non-obvious.

Claims 10, 12-14 and 23 are allowed.

Any comments considered necessary by applicant must be submitted no later than the payment of the issue fee and, to avoid processing delays, should preferably accompany the issue fee. Such submissions should be clearly labeled "Comments on Statement of Reasons for Allowance."

Application/Control Number: 10/631,958 Page 4

Art Unit: 1653

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Maryam Monshipouri whose telephone number is (571) 272-0932. The examiner can normally be reached on 7:00 a.m to 4:30 p.m. except for alternate Mondays.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Weber Jon P. can be reached on (571) 272-0925. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

Maryam Monshipouri Ph.D.

Primary Examiner

Title: Perfect score:

Sequence:

protein

Run on:

Scoring table:

Searched:

Minimum DB e Maximum DB e

Database :

ક્_{ટુ}

```
Sequence 11, Application US/09784810A

Sequence 11, Application US/09784810A

Patent No. US20020082203A1

GENERAL INFORMATION:

APPLICANT: RASTELLI, LUCA

TITLE OF INVENTION: NOVEL SPHINGOSINE KINASES AND NUCLEIC ACIDS ENCODING

FILE REFERENCE: 10716-08

CURRENT APPLICATION NUMBER: US/09/784,810A

CURRENT PILING DATE: 2001-02-14

PRIOR APPLICATION NUMBER: 60/191,261

PRIOR APPLICATION NUMBER: 60/191,261

PRIOR APPLICATION NUMBER: 60/191,261

SPRIOR APPLICATION NUMBER: 60/191,261

NUMBER OF SEQ ID NOS: 29

SOFTWARE: PATENTIN VEY: 2.1
                                                                                                                 Sequence 190238, Sequence 190238, Sequence 195284, Sequence 213665, Sequence 249555, Sequence 282512, Sequence 197631, Sequence 197631, Sequence 14927, A Sequence 199789, Sequence 19, Appl Sequence 19, Appl Sequence 19, Appl
                                                                                                                                                                                                                                                                                                                                                                        Sequence 19, Appl
Sequence 19, Appl
Sequence 28, Appl
Sequence 28, Appl
Sequence 8, Appli
Sequence 20, Appli
Sequence 20, Appli
Sequence 20, Appli
   Sequence 11, Appl
Sequence 2, Appli
Sequence 361076,
Sequence 29, Appl
                                                                    Sequence 29, Appl
Sequence 270874,
Sequence 133133,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20, Appl
Sequence 29, Appl
Sequence 29, Appl
Sequence 161184,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PKHLLVFINPFGGKGQGKRIYERKVAPLFTLASITTDIIGNKFYVNYVEVITEHANQAKE 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PKHLLVFINPFGGKGGGKRIYERKVAPLFTLASITTDIIGNKPYVNYVEVITEHANQAKE 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 1717; DB 9; Length 326; Best Local Similarity 100.0%; Pred. No. 2.8e-172; Matches 326; Conservative 0; Mismatches 0; Indels 0
6 US-10-631-958-11

4 US-10-315-597A-2

6 US-10-425-115-361076

US-10-425-115-37037

6 US-10-425-115-270874

6 US-10-425-115-270874

6 US-10-425-115-249553

6 US-10-425-115-249553

6 US-10-425-115-249555

6 US-10-425-115-26087

6 US-10-623-197631

6 US-10-623-185-288

6 US-10-623-185-288

6 US-10-623-201-19

6 US-10-623-201-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-876-281-8
US-10-053-510-20
US-10-052-510-20
US-10-622-011-20
US-10-348-052-29
US-10-622-011-29
US-10-632-011-29
US-10-6437-963-161184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ALIGNMENTS
     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                     RESULT 1
US-09-784-810A-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-784-810A-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             326
   1640.5
1632.5
                                   529
476.5
476.5
452
                                                                                                                                                                      321.5
318
311.5
305.5
305.5
305.5
300
293
                                                                                                                                                                                                                                                                                                                       288
287.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            270.5
270.5
270.5
270.5
270.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 11
       셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Appl
Sequence 1, Appl
Sequence 6, Appli
Sequence 10, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 11, Appl
Sequence 11, Appl
                                                                                                                                     (without alignments)
1990.064 Million cell updates/sec
                                                                                                                   ; Search time 64.5137 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 11,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2,
                                                                                                                                                                                                                     1 PKHLLVFINPFGGKGQGKRI......KCSRFNFLRFLIRHTNQQDQ 326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US11B_PUBCOMB.ppp:+
| cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.ppp:+
                     GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-784-810A-11
                                                                                                                                                                                                                                                                                                                                     Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                       1774312 segs, 393823214 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUMMARIES
                                                                                                                     September 3, 2005, 04:08:02
                                                                                                                                                                                                                                                                                                                                                                                                                       Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Published Applications AA:*
                                                                                     - protein search, using sw model
                                                                                                                                                                                                                                                     BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                          seq length: 0
seq length: 200000000
                                                                                                                                                                                    US-10-631-958-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query
Match 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0
100.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0
100.0
100.0
95.5
95.5
95.5
95.5
```

61 TLYEINIDKYDGIVCVGGDGMFSEVLHGLIGRTQRSAGVDQNHPRAVLVPSSLRIGIIPA 120

BEST AVAILABLE COPY

ઠે

1717 1640.5 1640.5 1640.5 1640.5

1717

Result

Gapa

```
181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a
a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 TLYEINIDKYDGIVCVGGDGMFSEVLHGLIGRTQRSAGVDQNHPRAVLVPSSLRIGIIPA 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       121 GSTDCVCYSTVGTSDAETSALHIVVGDSLAMDVSSVHHNSTLLRYSVSLLGYGFYGDIIK 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             181 DSEKKRMLGLARYDFSG/KTFLSHHCYEGTVSFLPAQHTVGSPRDRKPCRAGCFVCRQSK 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        241 QQLEBEQKKALYGLFAAEDVEEWQVVCGKFLAINATNMSCACRRSPRGLSPAAHLGDGSS 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 TLYBINIDKYDGIVCVGGDGMFSEVLHGLIGRTQRSAGVDQNHPRAVLVPSSLRIGIIPA 120
                                                               DSEKKRWIGLARYDFSGIKTFLSHHCYEGTVSFLPAQHTVGSPRDRKPCRAGCFVCRQSK 240
                                                                                          QQLEBEQKKALYGLEAAEDVEEWQVVCGKFLAINATWMSCACRRSPRGLSPAAHLGDGSS 300
                                                                                                                                                           241 QQLEEEQKKALYGLEAAEDVEEWQVVCGKFLAINATNMSCACRRSPRGLSPAAHLGDGSS 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 PKHILVFINPFGGKGGKRIYERKVAPLFTLASITTDIIGNKFYVNYVEVITEHANQAKE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 PKHLLVFINPFGGKGQGKRIYÉRKVAPLFTLASITTDIIGNKFYVNYVEVITEHANQAKE
             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.0%; Pred. No. 2.8e-172;
Matches 326; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/0996986;
Publication No. US20030125533A1
GEGRERAL INFORMATION:
GERREAL INFORMATION:
GERREAL INFORMATION:
APPLICANT: Sobbia
TITLE OF INVENTION: Regulation of human Sphingosine
TITLE OF INVENTION: Kinase-Like Protein
FILE REFERENCE: 004974.00594
CURRENT FILING DATE: 2001-10-04
PRIOR APPLICATION NUMBER: US 60/238,005
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2001-10-06
PRIOR FILING DATE: 2001-06
PRIOR FILING DATE: 2001-06
PRIOR FILING DATE: 2001-06
PRIOR FILING DATE: 2001-06
FROM REAL OF SECTION NUMBER: US 60/314,113
PRIOR FILING DATE: 2001-06
FROM REAL OF SECTION NUMBER: US 60/314,113
FROM REAL OF SECTION NOMBER: US 60/314,113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LRFLIRHTNOODO 326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DLILIRKCSRFNFLRFLIRHTNOODO 326
                                                                                                                                                                                                                           DLILIRKCSRFNFLRFLIRHTNQQDQ 326
                                                                                                                                                                                                                                                DLILIRKCSRFNFLRFLIRHTNQQDQ 326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 3
US-10-631-958-2
; Sequence 2, Application US/10631958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              301 DLILIRKCSRFNF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-969-896-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 2
                                                                                                            181
                                                                                                                                                                                                                             301
                                   121
                                                                                                                                                  241
                                                                         181
121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                유
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Š
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ઠ
                                                                                                          셤
                                                                                                                                                                               g
                                                                                                                                                                                                                           ò
                                                                         ò
                                                                                                                                                    ઢ
```

```
GENERAL INFOGRATION: NOVEL SPHINGO INE KINASES AND NUCLEIC ACIDS ENCODING TITLE OF INVENTION: NOVEL SPHINGO INE KINASES AND NUCLEIC ACIDS ENCODING TITLE OF INVENTION: SAME FILE REPERENCE: 10716-08

CURRENT APPLICATION NUMBER: US/09/P84,810

PRIOR FILING DATE: 2004-06-24

PRIOR FILING DATE: 2001-02-14

PRIOR FILING DATE: 2000-03-12

PRIOR APPLICATION NUMBER: 60/194,261

PRIOR APPLICATION NUMBER: 60/194,261

PRIOR FILING DATE: 2000-03-22

NUMBER OF SEQ ID NOS: 29

SOFTWARE: PATENTIN VET: 2.1

SEQ ID NO 11

SEQ ID NO 11

SEQ ID NO 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DSEKKRWIGLARYDPSGLKTPLSHHCYEGTVSFLPAQHTVGSPRDRKPCRAGCFVCRQSK 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               241 QOLEBEÇKKALYGLEAABDVBEWQVVÇKFLAINATNWSCACRRSPRGLSPAAHLGDGSS 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 TLYEINIDKYDGIVCVGGDGMFSEVLHGLIGFTQRSAGVDQNHPRAVLVPSSLRIGIIPA 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121 GSTDCVCXSTVGTSDARTSALHIVVGDSLAMDVSSVHHNSTLLRYSVSLLGYGFYGDIIK 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KFLAINATNMSCACRRSPRGLSPAAHLGDGSS 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 PKHLLVPINPFGGKGQGKRIYERKVAPLFTLASÍTTDIIGNKFYVNYVBVITEHANQAKB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 TLYBINIDKYDGIVÇVGGDGMFSEVLHGLIGRÍQRSAGVDQNHPRAVLVPSSLRIGIIPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 326;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indele
                                        APPLICANT: COSSIGA, Sophia
APPLICANT: ROSSIGA, Sophia
TITLE OF INVENTION: Requilation of human Sphingosine
TITLE OF INVENTION: Kinase-Like Protein
FILE REFERENCE: 004974.00594
CURRENT APPLICATION NUMBER: US/10/631,958
CURRENT FILING DATE: 2003-08-01
PRIOR FILING DATE: 2001-10-04
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: US 60/238,005
PRIOR FILING DATE: 2001-06
PRIOR APPLICATION NUMBER: US 60/314,113
PRIOR FILING DATE: 2010-10-06
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FRANKONS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 1717; DB 16;
100.0%; Pred. No. 2.84-172;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              QQLEEEQKKALYGLEAAEDVEEWQVVC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DLILIRKCSRFNFLRFLIRHTNQQDQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DLILIRKCSRFNFLRFLIRHTNQQDØ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 4
US-10-876-281-11
'Sequence 11, Application US/10876281
'Publication No. US20050123942A1
'GENERAL INFORMATION:
US20040192580A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 326; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 2
LENGTH: 326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-631-958-2
```

AHachmen

à g G ઠ g ઠ 셤 요 ò d ઠે g 8 ઠે ILIRKCSRPNFLRFLIRHTWQQDQFDFTFVEVYRVKKFQFTSKHMEDEDSDLKEGGKKRF 480 360 EKKRWLGLARYDFSGLKTFLSHHCYEGTVSFLPAQHTVGSPRDRKPCRAGCFVCRQSKQO 385 LEBEQKKALYGLEAAEDVEEWQVVCGKFLAINATNMSCACRRSPRGLSPAAHLGDGSSDL 420 266 TDCVCYSTVGTSDAETSALHIVVGDSLAMDVSSVHHNSTLLRYSVSLLGYGFYGDIIKDS 325 EMLEXLISRPKHLLVFINPFGGKGGCKRTYFERKVAPLFTLASITTDIIVTEHANQAKETL 180 206 YEINIDKYDGIVCVGGDGMFSEVLHGLIGHTUNGHTHTHTHTHTHTHTHTHTTTPAGS 265 300 145 82 GHICSSHPSCCCTVSNSS/NCDGEVLHSPAIEVRVHCQLVRLFARGIEENPKPDSHS 537 241 TDCVCYSTYGTSDARTSALHIVVGĎSLAMDVSSVHINSTLLRYSVSLLGYGFYGDIIKDS BKKRWLGLARYDFSGLKTPLSHÄCYEGTVSFLPAQHTVGSPRDRKPCRAGCFVCRQSKQQ 26 MGATGAABPLQSVLWVKQQRCAVSLEPARALJÆWWRSPGPGAGAPGADACSVPVSEIIAV **EETDVHGKHQGSGKWQKMEKPYAFTVHCVKKARRHRWKWAQVTFWCPEEQLCHLWLQTLR** 86 BETDVHGKHQGSGKWQKMEKPYAFTVHCVKRARRHRWKWAQVTFWCPEEQLCHLWLQTLR YBINIDKYDGIVCVGGDGMFSEVLHGLIGRIQRSAGVDQNHPRAVLVPSSLRIGIIPAGS Gaps Indels ö Mismatches . 537; Conservative US-10-631-958-11 301 326 361 386 481 206 181 421 61 121 Matches ò 셤 셤 ઠે 용 ઠે ઠે ઠે 셤 Š 셤 ઠે 셤 ò ద Š

26 MGATGAAEPLQSVIWVKQQRCAVSLEPARALLKWWRSPGPGAGAPGADACSVPVSEIIAV 85 0; Gaps 1 MGATGAARPIĄSYLWVKQQRCAVSLEPARALLRWWRSPGPGAGAPGADACSVPVSEIIAV Query Match
Best Local Similarity 100.0%; Score 2888; DB 16; Length 562;
Best Local Similarity 100.0%; Pred. No. 3.2e-278;
Matches 537; Conservative 0; Mismatches 0; Indels 0; GENERAL INFORMATION:
GENERAL INFORMATION:
FOR INVENTION: Regulation of human Sphingosine
TITLE OF INVENTION: Regulation of human Sphingosine
TITLE OF INVENTION: Ridade-Like Protein
FILE REFERENCE: 004974.00594
CURRENT APPLICATION NUMBER: US/10/631,958
CURRENT FILING DATE: 2001-10.04
PRIOR FILING DATE: 2001-10.04
PRIOR FILING DATE: 2001-10.04
PRIOR FILING DATE: 2001-10.06
PRIOR FILING DATE: 2001-10.06
PRIOR APPLICATION NUMBER: US 60/314,113
PRIOR APPLICATION NUMBER: US 60/314,113
PRIOR PILING DATE: 2001,08-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO II. Sequence 11, Application US/10631958 Publication No. US20040192580A1 GENERAL INFORMATION: ; TYPE: PRT ; ORGANISM: Homo sapien US-10-631-958-11

ઠ 음 ò 셤 ò 셤 ઠે

Query Match

99.7%; Score 2880; DB 14; Length 537;
Best Local Similarity 99.6%; Pred. No. 1.9e-277;
Matches 535; Conservative 1; Mismatches 1; Indels 0 Sequence 2, Application US/10315597A
Sequence 2, Application US/10315597A
Publication No. US20030162206A1
GENERAL INFORMATION:
APPLICANT: Sugiura, Masako
APPLICANT: Kohama, Takafumi
TITLE OF INVENTION: Cearamide Kinase and DNA Encoding It
TILE REFERENCE: O2658CIP/HG
CURRENT FILING DATE: 2002-12-10
PRIOR PLICATION NUMBER: US/10/315,597A
CURRENT FILING DATE: 2002-12-10
PRIOR PLICATION NUMBER: UP 2000-178039
PRIOR PLICATION NUMBER: UP 2000-178039
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Vo-SOFTWARE: Patentin Vo--ORGANISM: Homo sapiens US-10-315-597A-2 BESULT 7 US-10-315-597A-2 SEQ ID NO 2

61 BETDVHGKHQGSGKWQKMEKPYAPTVHCVKRARRHRWKWAQVTFWCPEGLCHLWLQTLR 120 EMLEKLTSRPKHLLVFINPFGGKGQGKRIYERKVAPLFTLASITTDIIVTEHANQAKETL 1 MGATGAAEPLOSVLWVKQQRCAVSLEPARALLRWWRSPGPGAGAPGADACSVPVSELIAV 181 YEINIDKYDGIVCVGGDGMFSEVLHGLIGRTQRSAGVDQNHPRAVLVPSSLRIGIIPAGS .. 0 121

ö

9

180

9

61 BETDVHGKHQGSGKWQKMEKPYAFTVHCVKRARRHRWKWAQVTFWCPBEQLCHLWLQTLR 120

BEST AVAILABLE COPY

```
TDCVCYSTVGTSDAETSALHIVVGDSLAMDVSSVHHNSTLLRYSVSLLGYGPYGDIIKDS 300
                                                                                  ILIRKCSKFWFLRFLIRHTWQDQFDFTFVEVYRVKKFQFTSKHMEDEDSDLKEGGKKRP 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STVGTSDAETSALHIVVGDSLAMDVSSVHHNSTLLRYSVSLLGYGFYGDIIKDSEKKRWL 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KALYGLEAAEDVEEWQVVCGKFLAINATNMSCACRESPRGLSPAAHLGDGSSDLILIRKC 360
                                                             EKKRWIGLARYDFSGLKTFLSHHCYEGTVSFLPAQHTVGSPRDRKPCRAGCFVCRQSKQQ 360
                                                                                                                             LEEEQKKALYGLEAAEDVEEWQVVCGKFLAINATWMSCACRRSPRGLSPAAHLGDGSSDL 420
                                                                                                                                                                                            ILIRKCSRFNFLRFLIRHTNQQDQFDFTFVEVYVKKFQFTSKHMEDEDSDLKEGGKKRF 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -----VTEHANQAKETLYEINID 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STVGTSDAETSALHIVVGDSLAMDVSSVHHNSTLLRYSVSLLGYGFYGDIIKDSEKKRWL 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: RASTELLY, LUCA
TITLE OF INVENTION: NOVEL SPHINGOSINE KINASES AND NUCLEIC ACIDS ENCODING
TITLE OF INVENTION: SAME
FILE OF INVENTION: SAME
FILE OF INVENTION: SAME
FILE OF INVENTION: SOURCE: 10716-08
CURRENT FILING DATE: 2001-02-14
PRIOR PILING DATE: 2001-02-14
PRIOR FILING DATE: 2000-02-14
PRIOR FILING DATE: 2000-03-22
NUMBER OF SEQ ID NOS: 29
SOUTWARE PATENTING DATE: 2000-03-22
NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                             537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 MEKPYAFIVHCVKRARRHRWKWAQVIFWCPEEQLCHLWLQTLREMLEKLJSRPKHLLVFI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KALYGLEAAEDVBEWQVVCGKFLAINATNMSCACRRSPRGLSPAAHLGDGSSDLILIRKC
TDCVCYSTVGTSDAETSALHIVVGDSLAMDVSSVHHNSTLLRYSVSLLGYGFYGDIIKDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHIMLOTLREMLEKLTSRPKHLLVFI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KYDGIVCVGGDGMFSEVLHGLIGRIQRSAGVDQNHPRAVLVPSSLRIGIIPAGSTDCVCY
                                                                                                                                                                                                                                                            GHICSSHPSCCCTVSNSSWNCDGEVLHSPAIEVRVHCQLVRLFARGIEENPKPDSHS
                                                                                                                                                                                                                                                                             GHICSSHPSCCCTVSNSSWNCDGEVLHSPAIEVRVHCQLVRLFARGIEENPKPDSHS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 9; Length 471;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 85.1%; Score 2456.5; DB 9
Best Local Similarity 97.5%; Pred. No. 2.44-235;
Matches 459; Conservative 1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78 MEKPYAFTVHCVKRARRHRWKWAQVTFWCPEEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NPFGGKGGGKRIYERKVAPLFTLASITTDII
                                                                                                                                                                                                                                                                                                                                                                           Sequence 6, Application US/09784810A
Patent No. US20020082203A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                             US-09-784-810A-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-784-810A-6
                                                                                          301
                                                                                                                                                                                                                         421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   301
241
                              241
                                                            301
                                                                                                                           361
                                                                                                                                                                                          421
                                                                                                                                                                                                                                                            481
                                                                                                                                                                                                                                                                                          481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ద
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8
                           g
                                                                                  a
                                                                                                                                                                                                                    g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ò
                                                            ò
                                                                                                                        ઠે
                                                                                                                                                   셤
                                                                                                                                                                                         ઠે
                                                                                                                                                                                                                                                          ઠે
                                                                                                                                                                                                                                                                                         원
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ò
```

```
246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 NPFGGKGGGKRIYERKVAPLFTLASITTDIIGNKFYVNYVEVITEHANQAKETLYEINID 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           181 STVGTSDAETSALHIVVGDSLAMDVSSVHHNSTLLRYSVSLLGYGFYGDIIKDSEKKRWL 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 366
486
                                   361 SRFNFLRFLIRHTNQQDQFDFTFVEVYRVKKFQFTGKHMEDEDSDLKEGGKKRFGHICSS 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    241 GLARYDFSGLKTFLSHHCYEGTVSFLPAQHTVGSPRDRKPCRAGCFVCROSKQQLEEEQX 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    420
                                                                                                                                                                                                                                                                                                KINASES AND NUCLEIC ACIDS ENCODING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 MEKPYAFTVHCVKRARRHRNKWAQVTFWCPEEQLCHLWLQTLREMLEKLISRPKHLLVFI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  138 NPFGGKGQGKRIYERKVAPLFTLASITTDII------VTEHANQAKETLYBINID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     301 KALYGLEAAEDVEEWQVVCGKFLAINATNMSCACRRSPRGLSPAHLGDGSSDLILIRKC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     78 MEKPYAFTVHCVKRARRHRWKWAQVTFWCPEEQLCHLWLQTLREMLEKLTSRPKHLLVFI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEVLHGLIGRIQRSAGVDQNHPRAVLVPSSLRIGIIPAGSTDCVCY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STVGTSDAETSALHIVVGDSLAMDVSSVHHNSTLLRYSVSLLGYGFYGDIIKDSEKKRWL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 307 GLARYDFSGLKTFLSHHCYEGTVSFLPAQHTVGSPRDRKPCRAGCFVCRQSKQQLEEEQK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              367 KALYGLEAAEDVEEWQVVCGKFLAINATNMSCACRRSPRGLSPAAHLGDGSSDLILIRKC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            427 SRPNFLRFLIRHTNQQDQFDFTFVEVYRVKKFQFTSKHMEDEDSDLKEGGKKRFGHICSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 361 SRFWFLRFLIRHTWQQDQPDFTFVEVYRVKKFQFTSKHMEDEDSDLKEGGKKRFGHICSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11; Gaps
                                                                                                421 HPSCCCTVSNSSWNCDGEVLHSPAIEVRVHCQLVRLFARGIEENPKPDSHS 471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TVSNSSWNCDGEVLHSPAIEVRVHCQLVRLFARGIEENPKPDSHS 537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 18; Length 471;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  421 HPSCCCTVSNSSWNCDGEVLHSPAIEVRVHCQLVRLFARGIEENPKPDSHS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 / Score 2456.5; DB 18
Pred. No. 2.4e-235;
1; Mismatches 0;
                                                                                                                                                                                                         Sequence 6, Application US/10876281

Publication No. US20050123942A1

GENERAL INPORMATION:

TITLE OF INVENTION: NOVEL SPHINGOSINE KINASES

TITLE OF INVENTION: NOWEL SPHINGOSINE KINASES

TITLE OF INVENTION: SAME

FILE REFERENCE: 10716-08

CURRENT APPLICATION NUMBER: US/10/876/281

CURRENT FILING DATE: 2004-06-24

PRIOR APPLICATION NUMBER: 60/182,360

PRIOR APPLICATION NUMBER: 60/182,360

PRIOR APPLICATION NUMBER: 60/191,261

PRIOR APPLICATION NUMBER: 60/191,261

PRIOR FILING DATE: 2000-03-22

NUMBER OF SEQ ID NOS: 29

SOFTWARE: PALENTIN VEY: 2.1

LENGTH: 471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 97.5%;
Matches 459; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           487 HPSCCC
                                                                                                                                                                           RESULT 9
US-10-876-281-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               247
                                                                             487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
                               g
                                                                                                              셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     쉽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ద
                                                                         ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8
```

RESULT 10 US-09-784-810A-11

Oy 446 ILIRKCSRENFLRFLIRHTNOODOFF DD 421 ILIRKCSRENFLRFLIRHTNOODOFF OY 506 GHICSSHPSCCCTVS4SSWNCDGEVI DD 481 GHICSSHPSCCGTVSNSSWNCDGEVI RESULT 12 US-10-315-697A-2 GEQUEOR 2) APPLICATION US/10315597A	ANT: Sugiura, Masako ANT: Kono, Keita ANT: Kono, Keita ANT: Kono, Keita ANT: Konoma, Takatumi OF INVENTION: Ceramide Kina, EFERENCE: 02658CIP/HG EFERENCE: 02658CIP/HG T APPLICATION NUMBER: US/10, T FILING DATE: 2000-06-14 OF SEQ ID NOS: 4 OF SEQ ID NOS: 4 FE: PatentIN Ver. 2.0 NO 2 NO 2 NO 2 NO 2 SEQ ID NOS: 4 FST FRI Homo sapiens	Matches 535; Conservative 1; Miss Qy	Db 121 EMLEKLISRPRHILVPINPFGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	326 301 386 386	446 421 506 481
Qy 326 EKKRWIGLARYDPSGLKTFLSHHCYEGTVSFLPAQHTWGSPRDRKPCRAGCFVCRQSKQQ Db 301 EKKRWIGLARYDPSGLKTFLSHHCYEGTVSFLPAQHTWGSPRDRKPCRAGCFVCRQSKQQ Qy 386 LEEEQKKALYGLEAAEDVEBWQVVCGKFLAINATWASCACRRSPRGLSPAAHLGDGSSDL 445 Db 361 LEEEQKKALYGLEAAEDVEBWQVVCGKFLAINATWASCACRRSPRGLSPAAHLGDGSSDL 420 QY 446 ILLIRKCSRFNFLRFLIRHTNQQDQFDFTFVEVTKKKQPTSKHMEDEDSDLKEGGKKRF 505 Db 421 ILIRKCSRFNFLRFLIRHTNQQDQFDFTFVEVTKKKQPTSKHMEDEDSDLKEGGKKRF 480 QY 506 GHICSSHPSCCCTVSNSSWNCDGEVLHSPALKRQVKRPQRTSRGHERNPKRDSHS 562	HESULT 11 Sequence 10, Application US/10631958 GENERAL INFORMATION: APPLICANT: Kossida, Sophia TITLE OF INVENTION: Kinase-Like Protein TITLE OF INVENTION: Kinase-Like Protein TITLE OF INVENTION: Kinase-Like Protein CURRENT FILING DATE: 2003-09-01 PRIOR PILING DATE: 2001-10-04 PRIOR FILING DATE: 2000-10-06 PRIOR PILING DATE: 2000-10-06 PRIOR PILING DATE: 2000-10-06 PRIOR PRILING DATE: 2000-10-06 PRIOR PILING DATE: 2000-10-06	NUMBER OF SEQ ID NOS: 16 SOFTWARE: FastSEQ for Windows Version 4.0	26 MGATGAAEPLOSVLWVKOORCAVSLEPARALLRWWRSPCPGAGAPGADACSVPVSEIIAV 85	21 21 7	24 10CVCYSTVGTSDAFISALHIVVOSSLAMDVSSVHHNSTLIRYSVSLLGYGFYGDIIKDS 325 10CVCYSTVGTSDAFISALHIVVOSSLAMDVSSVHHNSTLIRYSVSLLGYGFYGDIIKDS 325 241 TDCVCYSTVGTSDAFISALHIVVODSLAMDVSSVHHNSTLIRYSVSLLGYGFYGDIIKDS 300 326 EKKRWIGLARYDFSGLKTFLSHHCYEGTVSFLPAQHTVGSPRDKRECRAGCFVCRSKKQ 385

PDFTFVEVYRVKKFOPTSKHMEDEDSDLKEGGKKRF 505 VHCVKRARRHEWKWAQVTFWCPEEQLCHLWLQTLR 145 GLIGRTORSAGYDONHPRAVLVPSSLRIGIIPAGS 265 SLAMDVSSVHHNSTLLRYSVSLLCYGFYGDIIKDS 325 3KPLAINATNMSCACRRSPRGLSPAAHLGDGSSDL 445 BEARALLERWRRSPGPCAGAPCADACSVPVSELIAV 85 PTFVEVYRVKKFOFTSKHMEDEDSDLKEGGKKRF 505 VLHSPAIEVRVHCQLVRLFARGIEENPKPDSHS 562 20030162206 HSPAIEVRVHCQLVRLFARGIEENPKPDSHS 562 0; Gaps 1 2880; DB 29; Length 537; No. 3.2e-278; Smatches 1; Indels 0 and DNA Encoding It 5,597A 8039